Predicted TM Segments						TO THE TANK	
PHD Domain		ORIGICE CDEESPLITP OR GET GILRE OF IXSDIRCOLD OF IXS	DATONIC CANCEL	The same of the sa	GWCETAT DEDORTA E WARD DEGREE WAS A COTTAME OF THE COTTAME O	CRUCKSE GTPEKPLYMP CUST 63 1KF 1HQE CLUQULKUSRXEVELT	Origona assenlitepokotesloviolekkolorinessi origisele Origone codespiitpokoteslurungaadiopiiks3dtredelg
Accession	Number	BAA91319 AAH02122	AAF36360 BAB25080	XP_055276	BAA91173	BAA25523	AAH03404 AAH25394
	Mammalian	MARCH-I	MARCH-II	MARCH.	MARCH-V	MARCH-VI	MARCH-VIII MARCH-VIII

FIGURE 1A

	Accession	PHD Domain	Predicted TM Segments
Virus	Number		
BHV4-IE1	EDBED9	Hai ErgsesI iduky Ergisda oyung irung sasatroek fa	
HVS-ORF12		ELQACDCPSRVHHTCLQSHIQCFKSS	
KSHV-K3		TOT EVEL GNERRANGET GELEN WARSCHSTÜLT ISRNTAGO 15	
KSHV-K5		TWI CREEVENEAGONG	
LSDV-unnamed		į	
MHV68-K3		-	
Myxoma-m153R		TOTERCACD IVRYSEA THE COUNT WHEN THE THE TOTE OF THE TOTE OF THE TOTER OF THE	
Shope Fibroma-gp153R		į	
Swinepox-C7L			
Yaba*5L			

FIGURE 1B

Predicted TM Segments			
PHD Domain			CRUÇRC AQPDR CRIÇRC SUP QU CUI CLRC DEDRRRA
Accession	Number	T39453 NP 49530 NP 509788 NP 495728 NP 495728 NP 495821 NP 495821 NP 495821 NP 45624 AAF52133 AAF5624 AAF52133	AAF47637 AAF50284 AAF46341 AAF46173 AAF45871 AAF47637
	Other	S. Cerenvisiae S. pombe C. elegans D. melanogaster D. melanogaster D. melanogaster D. melanogaster	D. melanogaster D. melanogaster D. melanogaster D. melanogaster D. melanogaster E. cuniculi

FIGURE 1C

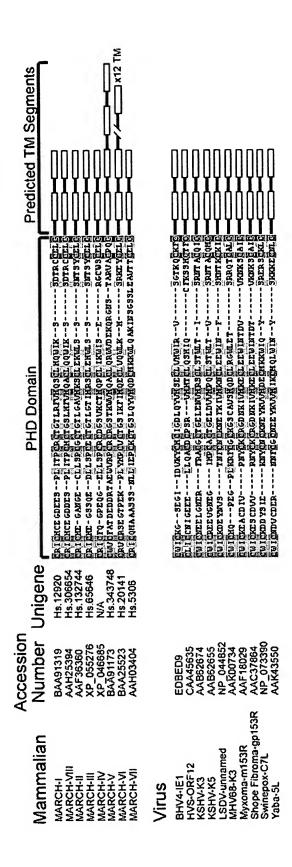


FIGURE 1D

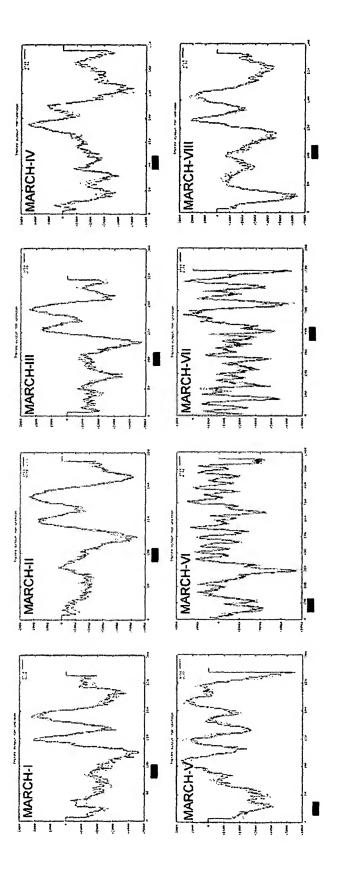
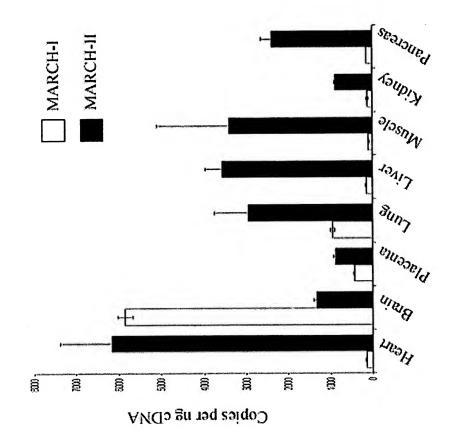


FIGURE 1E





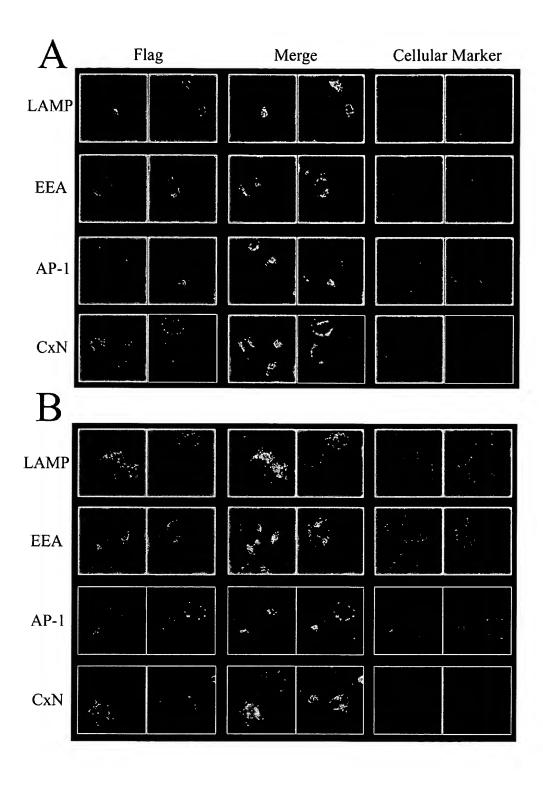
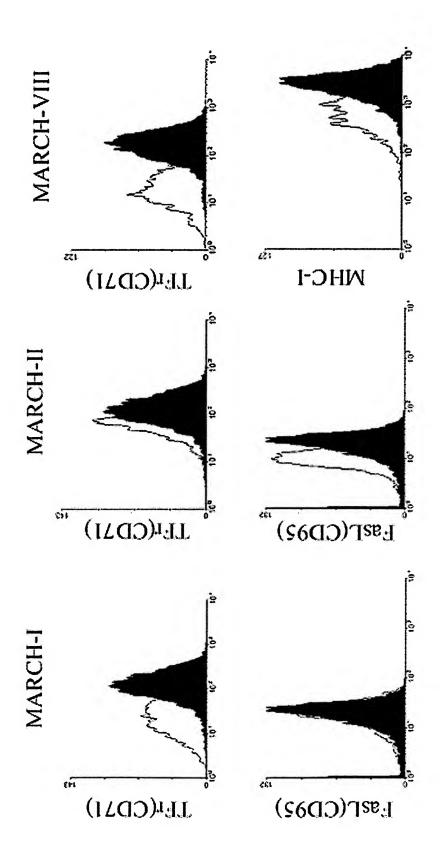


FIGURE 3



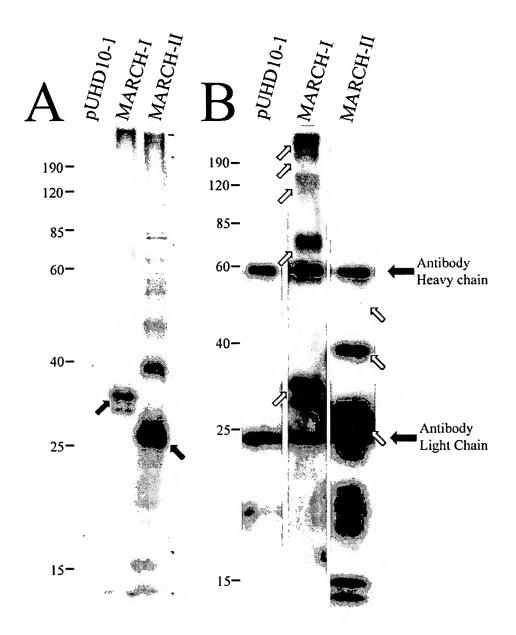


FIGURE 5

BEST AVAILABLE COPY

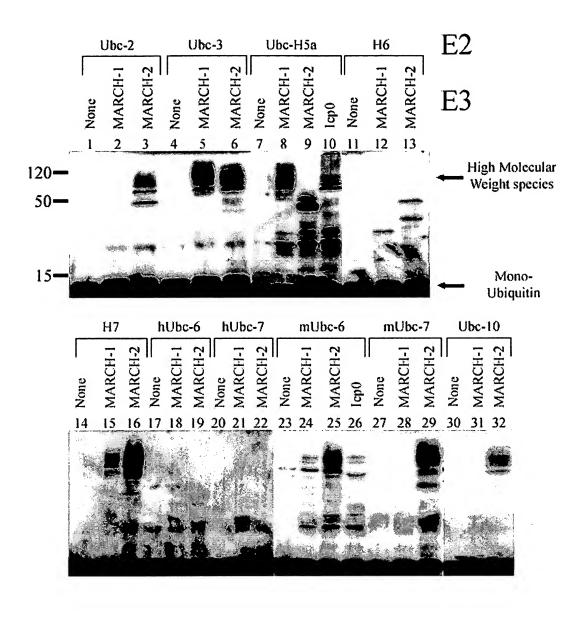


FIGURE 6

BEST AVAILABLE COPY

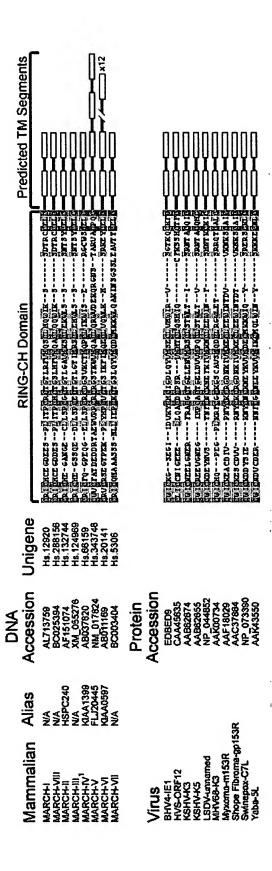


FIGURE 7A

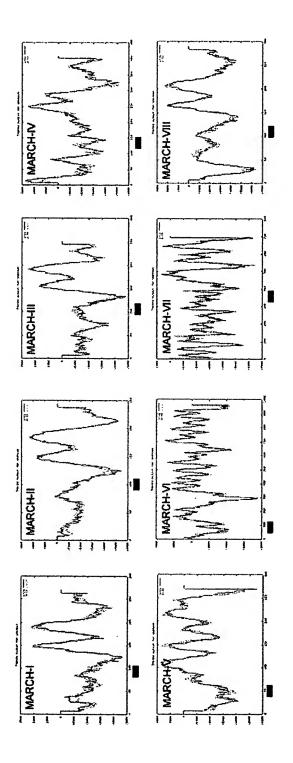
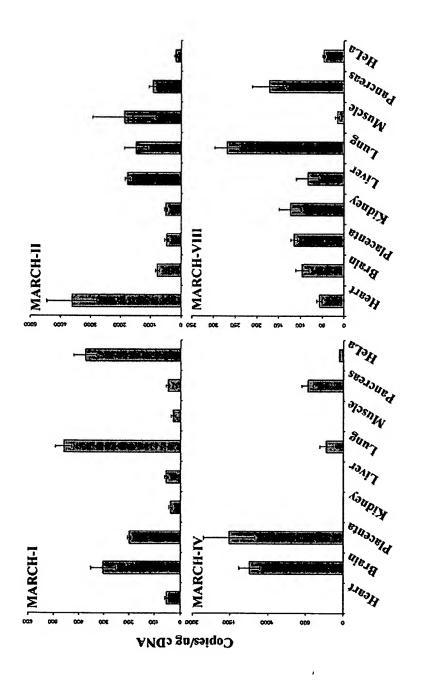


FIGURE 7B



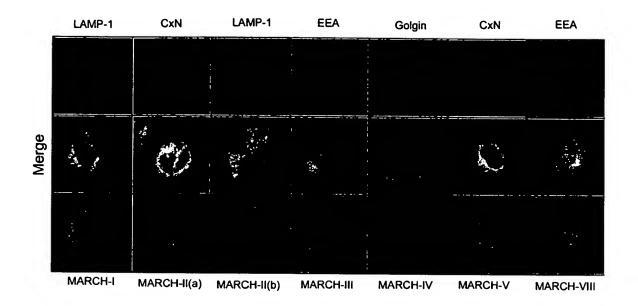


FIGURE 8

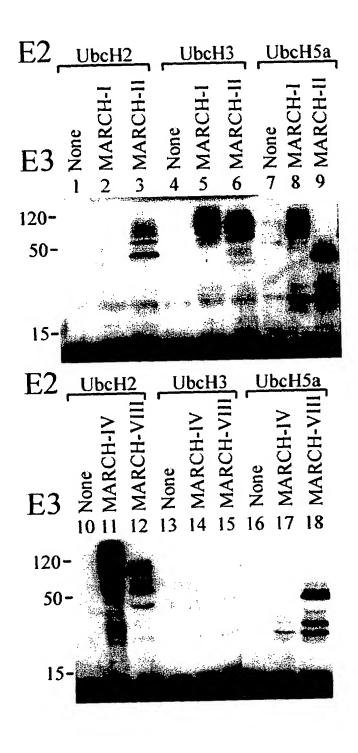


FIGURE 9

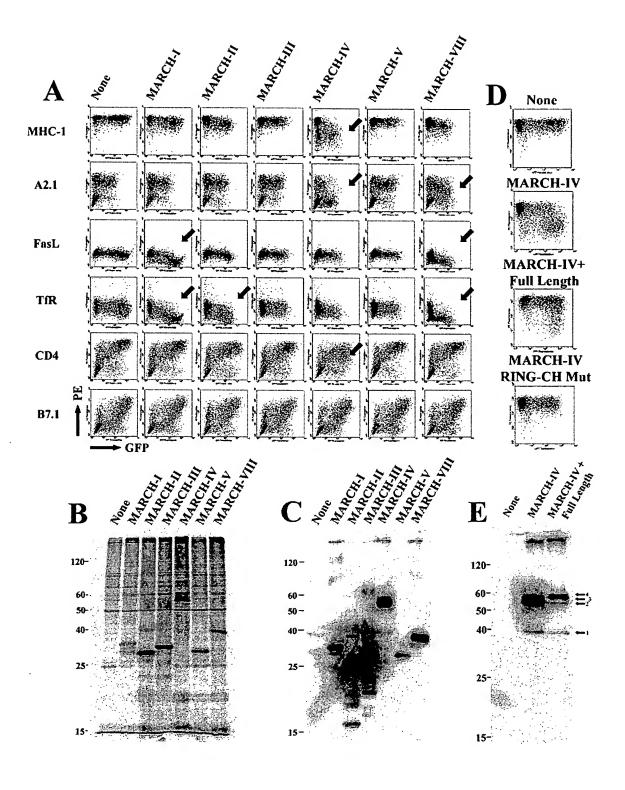


FIGURE 10

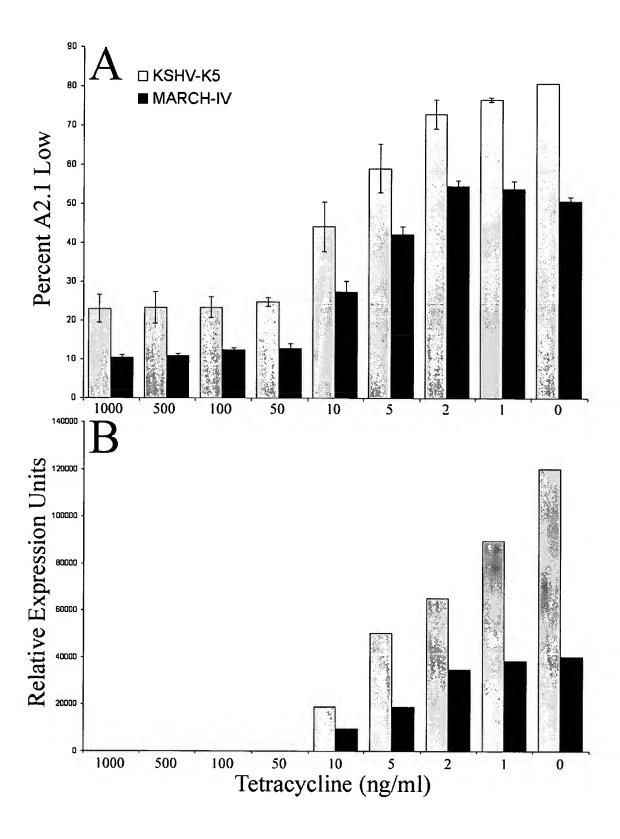


FIGURE 11

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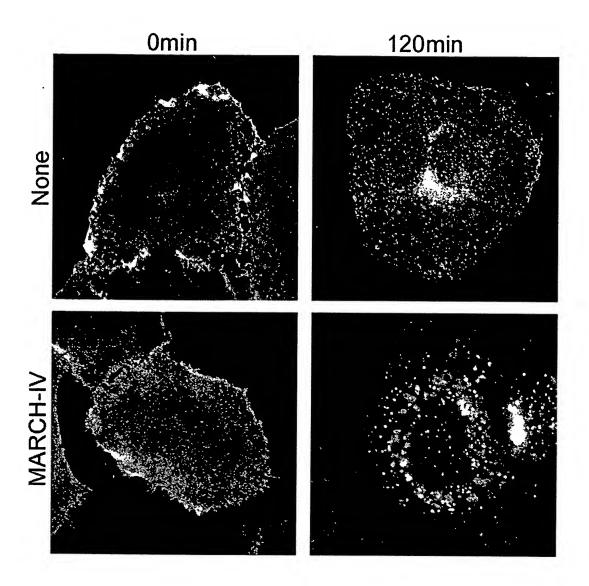


FIGURE 12A

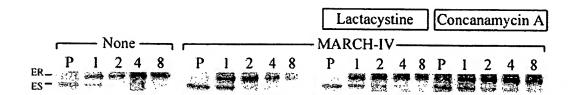


FIGURE 12B

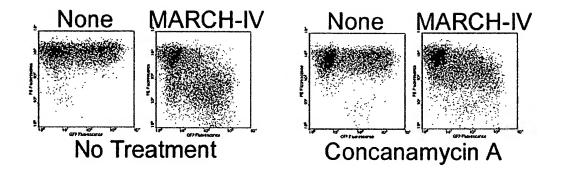


FIGURE 12C

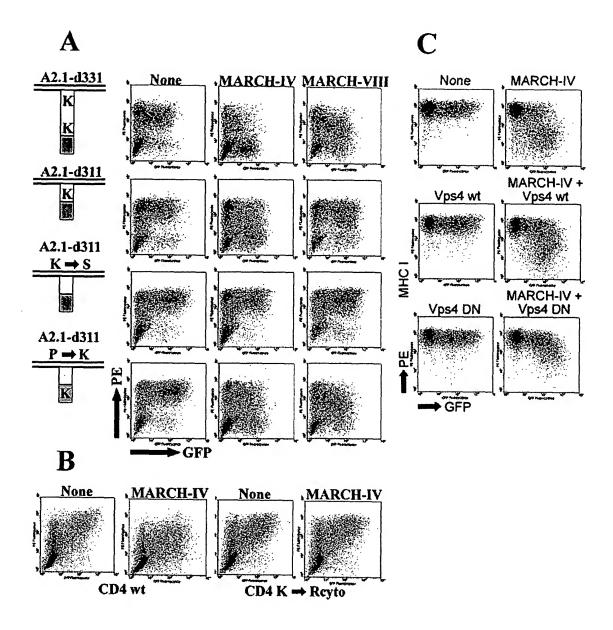


FIGURE 13